

SEQUENCE LISTING

<110> Boehringer Ingelheim Vetmedica GmbH

<120> Recombinant attenuation of Porcine Reproductive and Respiratory Syndrome (PRRSV)

<130> 1-1120,25,26-1-C1

<140> to be assigned

<141> June 20, 2003

<150> 09/772,316

<151> January 26, 2001

<150> DE 10003371.7

<151> 2000-01-26

<150> DE 10003373.3

<151> 2000-01-26

<150> DE 10003372.5

<151> 2000-01-26

<160> 6

<170> PatentIn Ver. 2.1

<210> 1

<211> 2502

<212> PRT

<213> Porcine reproductive and respiratory syndrome virus

<400> 1

```

Met Ser Gly Ile Leu Asp Arg Cys Thr Cys Thr Pro Asn Ala Arg Val
 1             5             10             15

Phe Met Ala Glu Gly Gln Val Tyr Cys Thr Arg Cys Leu Ser Ala Arg
      20             25             30

Ser Leu Leu Pro Leu Asn Leu Gln Val Ser Glu Leu Gly Val Leu Gly
      35             40             45

Leu Phe Tyr Arg Pro Glu Glu Pro Leu Arg Trp Thr Leu Pro Arg Ala
      50             55             60

Phe Pro Thr Val Glu Cys Ser Pro Ala Gly Ala Cys Trp Leu Ser Ala
      65             70             75             80

Ile Phe Pro Ile Ala Arg Met Thr Ser Gly Asn Leu Asn Phe Gln Gln
      85             90             95

Arg Met Val Arg Val Ala Ala Glu Leu Tyr Arg Ala Gly Gln Leu Thr
      100            105            110

Pro Ala Val Leu Lys Ala Leu Gln Val Tyr Glu Arg Gly Cys Arg Trp
      115            120            125

```

Tyr	Pro	Ile	Val	Gly	Pro	Val	Pro	Gly	Val	Ala	Val	Phe	Ala	Asn	Ser	130	135	140	
Leu	His	Val	Ser	Asp	Lys	Pro	Phe	Pro	Gly	Ala	Thr	His	Val	Leu	Thr	145	150	155	160
Asn	Leu	Pro	Leu	Pro	Gln	Arg	Pro	Lys	Pro	Glu	Asp	Phe	Cys	Pro	Phe	165	170	175	
Glu	Cys	Ala	Met	Ala	Thr	Val	Tyr	Asp	Ile	Gly	His	Asp	Ala	Val	Met	180	185	190	
Tyr	Val	Ala	Glu	Arg	Lys	Val	Ser	Trp	Ala	Pro	Arg	Gly	Gly	Asp	Glu	195	200	205	
Val	Lys	Phe	Glu	Ala	Val	Pro	Gly	Glu	Leu	Lys	Leu	Ile	Ala	Asn	Arg	210	215	220	
Leu	Arg	Thr	Ser	Phe	Pro	Pro	His	His	Thr	Val	Asp	Met	Ser	Lys	Phe	225	230	235	240
Ala	Phe	Thr	Ala	Pro	Gly	Cys	Gly	Val	Ser	Met	Arg	Val	Glu	Arg	Gln	245	250	255	
His	Gly	Cys	Leu	Pro	Ala	Asp	Thr	Val	Pro	Glu	Gly	Asn	Cys	Trp	Trp	260	265	270	
Ser	Leu	Phe	Asp	Leu	Leu	Pro	Leu	Glu	Val	Gln	Asn	Lys	Glu	Ile	Arg	275	280	285	
His	Ala	Asn	Gln	Phe	Gly	Tyr	Gln	Thr	Lys	His	Gly	Val	Ser	Gly	Lys	290	295	300	
Tyr	Leu	Gln	Arg	Arg	Leu	Gln	Val	Asn	Gly	Leu	Arg	Ala	Val	Thr	Asp	305	310	315	320
Leu	Asn	Gly	Pro	Ile	Val	Val	Gln	Tyr	Phe	Ser	Val	Lys	Glu	Ser	Trp	325	330	335	
Ile	Arg	His	Leu	Lys	Leu	Ala	Gly	Glu	Pro	Ser	Tyr	Ser	Gly	Phe	Glu	340	345	350	
Asp	Leu	Leu	Arg	Ile	Arg	Val	Glu	Pro	Asn	Thr	Ser	Pro	Leu	Ala	Asp	355	360	365	
Lys	Glu	Glu	Lys	Ile	Phe	Arg	Phe	Gly	Ser	His	Lys	Trp	Tyr	Gly	Ala	370	375	380	
Gly	Lys	Arg	Ala	Arg	Lys	Ala	Arg	Ser	Cys	Ala	Thr	Ala	Thr	Val	Ala	385	390	395	400
Gly	Arg	Ala	Leu	Ser	Val	Arg	Glu	Thr	Arg	Gln	Ala	Lys	Glu	His	Glu	405	410	415	
Val	Ala	Gly	Ala	Asn	Lys	Ala	Glu	His	Leu	Lys	His	Tyr	Ser	Pro	Pro	420	425	430	

Ala	Glu	Gly	Asn	Cys	Gly	Trp	His	Cys	Ile	Ser	Ala	Ile	Ala	Asn	Arg		
		435					440					445					
Met	Val	Asn	Ser	Lys	Phe	Glu	Thr	Thr	Leu	Pro	Glu	Arg	Val	Arg	Pro		
	450					455					460						
Pro	Asp	Asp	Trp	Ala	Thr	Asp	Glu	Asp	Leu	Val	Asn	Ala	Ile	Gln	Ile		
465					470					475					480		
Leu	Arg	Leu	Pro	Ala	Ala	Leu	Asp	Arg	Asn	Gly	Ala	Cys	Thr	Ser	Ala		
				485					490					495			
Lys	Tyr	Val	Leu	Lys	Leu	Glu	Gly	Glu	His	Trp	Thr	Val	Thr	Val	Thr		
		500						505					510				
Pro	Gly	Met	Ser	Pro	Ser	Leu	Leu	Pro	Leu	Glu	Cys	Val	Gln	Gly	Cys		
		515					520					525					
Cys	Gly	His	Lys	Gly	Gly	Leu	Gly	Ser	Pro	Asp	Ala	Val	Glu	Val	Ser		
	530					535					540						
Gly	Phe	Asp	Pro	Ala	Cys	Leu	Asp	Arg	Leu	Ala	Glu	Val	Met	His	Leu		
545					550					555					560		
Pro	Ser	Ser	Ala	Ile	Pro	Ala	Ala	Leu	Ala	Glu	Met	Ser	Gly	Asp	Ser		
				565					570					575			
Asp	Arg	Ser	Ala	Ser	Pro	Val	Thr	Thr	Val	Trp	Thr	Val	Ser	Gln	Phe		
			580					585					590				
Phe	Ala	Arg	His	Ser	Gly	Gly	Asn	His	Pro	Asp	Gln	Val	Arg	Leu	Gly		
		595					600					605					
Lys	Ile	Ile	Ser	Leu	Cys	Gln	Val	Ile	Glu	Asp	Cys	Cys	Cys	Ser	Gln		
	610					615					620						
Asn	Lys	Thr	Asn	Arg	Val	Thr	Pro	Glu	Glu	Val	Ala	Ala	Lys	Ile	Asp		
625					630					635					640		
Leu	Tyr	Leu	Arg	Gly	Ala	Thr	Asn	Leu	Glu	Glu	Cys	Leu	Ala	Arg	Leu		
				645					650					655			
Glu	Lys	Ala	Arg	Pro	Pro	Arg	Val	Ile	Asp	Thr	Ser	Phe	Asp	Trp	Asp		
			660					665					670				
Val	Val	Leu	Pro	Gly	Val	Glu	Ala	Ala	Thr	Gln	Thr	Ile	Lys	Leu	Pro		
		675					680					685					
Gln	Val	Asn	Gln	Cys	Arg	Ala	Leu	Val	Pro	Val	Val	Thr	Gln	Lys	Ser		
	690					695					700						
Leu	Asp	Asn	Asn	Ser	Val	Pro	Leu	Thr	Ala	Phe	Ser	Leu	Ala	Asn	Tyr		
705					710					715					720		
Tyr	Tyr	Arg	Ala	Gln	Gly	Asp	Glu	Val	Arg	His	Arg	Glu	Arg	Leu	Thr		
				725					730					735			

Ala	Val	Leu	Ser	Lys	Leu	Glu	Lys	Val	Val	Arg	Glu	Glu	Tyr	Gly	Leu	
			740					745					750			
Met	Pro	Thr	Glu	Pro	Gly	Pro	Arg	Pro	Thr	Leu	Pro	Arg	Gly	Leu	Asp	
		755					760					765				
Glu	Leu	Lys	Asp	Gln	Met	Glu	Glu	Asp	Leu	Leu	Lys	Leu	Ala	Asn	Ala	
	770					775					780					
Gln	Thr	Thr	Ser	Asp	Met	Met	Ala	Trp	Ala	Val	Glu	Gln	Val	Asp	Leu	
785					790					795					800	
Lys	Thr	Trp	Val	Lys	Asn	Tyr	Pro	Arg	Trp	Thr	Pro	Pro	Pro	Pro	Pro	
				805					810					815		
Pro	Lys	Val	Gln	Pro	Arg	Lys	Thr	Lys	Pro	Val	Lys	Ser	Leu	Pro	Glu	
			820					825					830			
Arg	Lys	Pro	Val	Pro	Ala	Pro	Arg	Arg	Lys	Val	Gly	Ser	Asp	Cys	Gly	
		835					840					845				
Ser	Pro	Val	Ser	Leu	Gly	Gly	Asp	Val	Pro	Asn	Ser	Trp	Glu	Asp	Leu	
	850					855					860					
Ala	Val	Ser	Ser	Pro	Phe	Asp	Leu	Pro	Thr	Pro	Pro	Glu	Pro	Ala	Thr	
865					870					875					880	
Pro	Ser	Ser	Glu	Leu	Val	Ile	Val	Ser	Ser	Pro	Gln	Cys	Ile	Phe	Arg	
				885					890					895		
Pro	Ala	Thr	Pro	Leu	Ser	Glu	Pro	Ala	Pro	Ile	Pro	Ala	Pro	Arg	Gly	
			900					905					910			
Thr	Val	Ser	Arg	Pro	Val	Thr	Pro	Leu	Ser	Glu	Pro	Ile	Pro	Val	Pro	
		915					920						925			
Ala	Pro	Arg	Arg	Lys	Phe	Gln	Gln	Val	Lys	Arg	Leu	Ser	Ser	Ala	Ala	
	930					935					940					
Ala	Ile	Pro	Pro	Tyr	Gln	Asp	Glu	Pro	Leu	Asp	Leu	Ser	Ala	Ser	Ser	
945					950					955					960	
Gln	Thr	Glu	Tyr	Glu	Ala	Ser	Pro	Pro	Ala	Pro	Pro	Gln	Ser	Gly	Gly	
				965					970					975		
Val	Leu	Gly	Val	Glu	Gly	His	Glu	Ala	Glu	Glu	Thr	Leu	Ser	Glu	Ile	
			980					985					990			
Ser	Asp	Met	Ser	Gly	Asn	Ile	Lys	Pro	Ala	Ser	Val	Ser	Ser	Ser	Ser	
		995				1000						1005				
Ser	Leu	Ser	Ser	Val	Arg	Ile	Thr	Arg	Pro	Lys	Tyr	Ser	Ala	Gln	Ala	
	1010					1015					1020					
Ile	Ile	Asp	Ser	Gly	Gly	Pro	Cys	Ser	Gly	His	Leu	Gln	Glu	Val	Lys	
1025					1030					1035					1040	

Glu Thr Cys Leu Ser Val Met Arg Glu Ala Cys Asp Ala Thr Lys Leu	1045	1050	1055
Asp Asp Pro Ala Thr Gln Glu Trp Leu Ser Arg Met Trp Asp Arg Val	1060	1065	1070
Asp Met Leu Thr Trp Arg Asn Thr Ser Val Tyr Gln Ala Ile Cys Thr	1075	1080	1085
Leu Asp Gly Arg Leu Lys Phe Leu Pro Lys Met Ile Leu Glu Thr Pro	1090	1095	1100
Pro Pro Tyr Pro Cys Glu Phe Val Met Met Pro His Thr Pro Ala Pro	1105	1110	1115
Ser Val Gly Ala Glu Ser Asp Leu Thr Ile Gly Ser Val Ala Thr Glu	1125	1130	1135
Asp Val Pro Arg Ile Leu Glu Lys Ile Glu Asn Val Gly Glu Met Ala	1140	1145	1150
Asn Gln Gly Pro Leu Ala Phe Ser Glu Asp Lys Pro Val Asp Asp Gln	1155	1160	1165
Leu Val Asn Asp Pro Arg Ile Ser Ser Arg Arg Pro Asp Glu Ser Thr	1170	1175	1180
Ser Ala Pro Ser Ala Gly Thr Gly Gly Ala Gly Ser Phe Thr Asp Leu	1185	1190	1195
Pro Pro Ser Asp Gly Ala Asp Ala Asp Gly Gly Gly Pro Phe Arg Thr	1205	1210	1215
Val Lys Arg Lys Ala Glu Arg Leu Phe Asp Gln Leu Ser Arg Gln Val	1220	1225	1230
Phe Asp Leu Val Ser His Leu Pro Val Phe Phe Ser Arg Leu Phe Tyr	1235	1240	1245
Pro Gly Gly Gly Tyr Ser Pro Gly Asp Trp Gly Phe Ala Ala Phe Thr	1250	1255	1260
Leu Leu Cys Leu Phe Leu Cys Tyr Ser Tyr Pro Ala Phe Gly Ile Ala	1265	1270	1275
Pro Leu Leu Gly Val Phe Ser Gly Ser Ser Arg Arg Val Arg Met Gly	1285	1290	1295
Val Phe Gly Cys Trp Leu Ala Phe Ala Val Gly Leu Phe Lys Pro Val	1300	1305	1310
Ser Asp Pro Val Gly Ala Ala Cys Glu Phe Asp Ser Pro Glu Cys Arg	1315	1320	1325
Asn Ile Leu His Ser Phe Glu Leu Leu Lys Pro Trp Asp Pro Val Arg	1330	1335	1340

Ser Leu Val Val Gly Pro Val Gly Leu Gly Leu Ala Ile Leu Gly Arg
 1345 1350 1355 1360
 Leu Leu Gly Gly Ala Arg Cys Ile Trp His Phe Leu Leu Arg Leu Gly
 1365 1370 1375
 Ile Val Ala Asp Cys Ile Leu Ala Gly Ala Tyr Val Leu Ser Gln Gly
 1380 1385 1390
 Arg Cys Lys Lys Cys Trp Gly Ser Cys Ile Arg Thr Ala Pro Asn Glu
 1395 1400 1405
 Val Ala Phe Asn Val Phe Pro Phe Thr Arg Ala Thr Arg Ser Ser Leu
 1410 1415 1420
 Ile Asp Leu Cys Asp Arg Phe Cys Ala Pro Lys Gly Met Asp Pro Ile
 1425 1430 1435 1440
 Phe Leu Ala Thr Gly Trp Arg Gly Cys Trp Ala Gly Arg Ser Pro Ile
 1445 1450 1455
 Glu Gln Pro Ser Glu Lys Pro Ile Ala Phe Ala Gln Leu Asp Glu Lys
 1460 1465 1470
 Lys Ile Thr Ala Arg Thr Val Val Ala Gln Pro Tyr Asp Pro Asn Gln
 1475 1480 1485
 Ala Val Lys Cys Leu Arg Val Leu Gln Ser Gly Gly Arg Trp Trp Leu
 1490 1495 1500
 Ser Gly Pro Lys Ser Gly Gln Gly Phe Arg Cys Ser Ile Pro Ser Pro
 1505 1510 1515 1520
 Phe Phe Pro Thr Gly Val Lys Val Asp Pro Asp Cys Arg Val Val Val
 1525 1530 1535
 Asp Pro Asp Thr Phe Thr Ala Ala Leu Arg Ser Gly Tyr Ser Thr Thr
 1540 1545 1550
 Asn Leu Val Leu Gly Val Gly Asp Phe Ala Gln Leu Asn Gly Leu Lys
 1555 1560 1565
 Ile Arg Gln Ile Ser Lys Pro Ser Gly Gly Gly Pro His Leu Met Ala
 1570 1575 1580
 Ala Leu His Val Ala Cys Ser Met Ala Leu His Met Leu Ala Gly Ile
 1585 1590 1595 1600
 Tyr Val Thr Ala Val Gly Ser Cys Gly Thr Gly Thr Asn Asp Pro Trp
 1605 1610 1615
 Cys Ala Asn Pro Phe Ala Val Pro Gly Tyr Gly Pro Gly Ser Leu Cys
 1620 1625 1630
 Thr Ser Arg Leu Cys Ile Ser Gln His Gly Leu Thr Leu Pro Leu Thr
 1635 1640 1645

Ala Leu Val Ala Gly Phe Gly Ile Gln Glu Ile Ala Leu Val Val Leu
 1650 1655 1660
 Ile Phe Val Ser Ile Gly Gly Met Ala His Arg Leu Ser Cys Lys Ala
 1665 1670 1675 1680
 Asp Met Leu Cys Val Leu Leu Ala Ile Ala Ser Tyr Val Trp Val Pro
 1685 1690 1695
 Leu Thr Trp Leu Leu Cys Val Phe Pro Cys Trp Leu Arg Cys Phe Ser
 1700 1705 1710
 Leu His Pro Leu Thr Ile Leu Trp Leu Val Phe Phe Leu Ile Ser Val
 1715 1720 1725
 Asn Met Pro Ser Gly Ile Leu Ala Met Val Leu Leu Val Ser Leu Trp
 1730 1735 1740
 Leu Leu Gly Arg Tyr Thr Asn Val Ala Gly Leu Val Thr Pro Tyr Asp
 1745 1750 1755 1760
 Ile His His Tyr Thr Ser Gly Pro Arg Gly Val Ala Ala Leu Ala Thr
 1765 1770 1775
 Ala Pro Asp Gly Thr Tyr Leu Ala Ala Val Arg Arg Ala Ala Leu Thr
 1780 1785 1790
 Gly Arg Thr Met Leu Phe Thr Pro Ser Gln Leu Gly Ser Leu Leu Glu
 1795 1800 1805
 Gly Ala Phe Arg Thr Arg Lys Pro Ser Leu Asn Thr Val Asn Val Ile
 1810 1815 1820
 Gly Ser Ser Met Gly Ser Gly Gly Val Phe Thr Ile Asp Gly Lys Val
 1825 1830 1835 1840
 Lys Cys Val Thr Ala Ala His Val Leu Thr Gly Asn Ser Ala Arg Val
 1845 1850 1855
 Ser Gly Val Gly Phe Asn Gln Met Leu Asp Phe Asp Val Lys Gly Asp
 1860 1865 1870
 Phe Ala Ile Ala Asp Cys Pro Asn Trp Gln Gly Ala Ala Pro Lys Thr
 1875 1880 1885
 Gln Phe Cys Thr Asp Gly Trp Thr Gly Arg Ala Tyr Trp Leu Thr Ser
 1890 1895 1900
 Ser Gly Val Glu Pro Gly Val Ile Gly Lys Gly Phe Ala Phe Cys Phe
 1905 1910 1915 1920
 Thr Ala Cys Gly Asp Ser Gly Ser Pro Val Ile Thr Glu Ala Gly Glu
 1925 1930 1935
 Leu Val Gly Val His Thr Gly Ser Asn Lys Gln Gly Gly Gly Ile Val
 1940 1945 1950

Thr	Arg	Pro	Ser	Gly	Gln	Phe	Cys	Asn	Val	Ala	Pro	Ile	Lys	Leu	Ser	1955	1960	1965	
Glu	Leu	Ser	Glu	Phe	Phe	Ala	Gly	Pro	Lys	Val	Pro	Leu	Gly	Asp	Val	1970	1975	1980	
Lys	Val	Gly	Ser	His	Ile	Ile	Lys	Asp	Ile	Ser	Glu	Val	Pro	Ser	Asp	1985	1990	1995	2000
Leu	Cys	Ala	Leu	Leu	Ala	Ala	Lys	Pro	Glu	Leu	Glu	Gly	Gly	Leu	Ser	2005	2010	2015	
Thr	Val	Gln	Leu	Leu	Cys	Val	Phe	Phe	Leu	Leu	Trp	Arg	Met	Met	Gly	2020	2025	2030	
His	Ala	Trp	Thr	Pro	Leu	Val	Ala	Val	Ser	Phe	Phe	Ile	Leu	Asn	Glu	2035	2040	2045	
Val	Leu	Pro	Ala	Val	Leu	Val	Arg	Ser	Val	Phe	Ser	Phe	Gly	Met	Phe	2050	2055	2060	
Val	Leu	Ser	Trp	Leu	Thr	Pro	Trp	Ser	Ala	Gln	Val	Leu	Met	Ile	Arg	2065	2070	2075	2080
Leu	Leu	Thr	Ala	Ala	Leu	Asn	Arg	Asn	Arg	Trp	Ser	Leu	Ala	Phe	Phe	2085	2090	2095	
Ser	Leu	Gly	Ala	Val	Thr	Gly	Phe	Val	Ala	Asp	Leu	Ala	Ala	Thr	Gln	2100	2105	2110	
Gly	His	Pro	Leu	Gln	Ala	Val	Met	Asn	Leu	Ser	Thr	Tyr	Ala	Phe	Leu	2115	2120	2125	
Pro	Arg	Met	Met	Val	Val	Thr	Ser	Pro	Val	Pro	Val	Ile	Thr	Cys	Gly	2130	2135	2140	
Val	Val	His	Leu	Leu	Ala	Ile	Ile	Leu	Tyr	Leu	Phe	Lys	Tyr	Arg	Gly	2145	2150	2155	2160
Pro	His	His	Ile	Leu	Val	Gly	Asp	Gly	Val	Phe	Ser	Ala	Ala	Phe	Phe	2165	2170	2175	
Leu	Arg	Tyr	Phe	Ala	Glu	Gly	Lys	Leu	Arg	Glu	Gly	Val	Ser	Gln	Ser	2180	2185	2190	
Cys	Gly	Met	Asn	His	Glu	Ser	Leu	Thr	Gly	Ala	Leu	Ala	Met	Arg	Leu	2195	2200	2205	
Asn	Asp	Glu	Asp	Leu	Asp	Phe	Leu	Met	Lys	Trp	Thr	Asp	Phe	Lys	Cys	2210	2215	2220	
Phe	Val	Ser	Ala	Ser	Asn	Met	Arg	Asn	Ala	Ala	Gly	Gln	Phe	Ile	Glu	2225	2230	2235	2240
Ala	Ala	Tyr	Ala	Lys	Ala	Leu	Arg	Val	Glu	Leu	Ala	Gln	Leu	Val	Gln	2245	2250	2255	

Val Asp Lys Val Arg Gly Thr Leu Ala Lys Leu Glu Ala Phe Ala Asp
 2260 2265 2270
 Thr Val Ala Pro Gln Leu Ser Pro Gly Asp Ile Val Val Ala Leu Gly
 2275 2280 2285
 His Thr Pro Val Gly Ser Ile Phe Asp Leu Lys Val Gly Ser Thr Lys
 2290 2295 2300
 His Thr Leu Gln Ala Ile Glu Thr Arg Val Leu Ala Gly Ser Lys Met
 2305 2310 2315 2320
 Thr Val Ala Arg Val Val Asp Pro Thr Pro Thr Pro Pro Ala Pro
 2325 2330 2335
 Val Pro Ile Pro Leu Pro Pro Lys Val Leu Glu Asn Gly Pro Asn Ala
 2340 2345 2350
 Trp Gly Asp Glu Asp Arg Leu Asn Lys Lys Lys Arg Arg Arg Met Glu
 2355 2360 2365
 Ala Leu Gly Ile Tyr Val Met Gly Gly Lys Lys Tyr Gln Lys Phe Trp
 2370 2375 2380
 Asp Lys Asn Ser Gly Asp Val Phe Tyr Glu Glu Val His Asn Asn Thr
 2385 2390 2395 2400
 Asp Glu Trp Glu Cys Leu Arg Val Gly Asp Pro Ala Asp Phe Asp Pro
 2405 2410 2415
 Glu Lys Gly Thr Leu Cys Gly His Val Thr Ile Glu Asn Lys Ala Tyr
 2420 2425 2430
 His Val Tyr Thr Ser Pro Ser Gly Lys Lys Phe Leu Val Pro Val Asn
 2435 2440 2445
 Pro Glu Asn Gly Arg Val Gln Trp Glu Ala Ala Lys Leu Ser Val Glu
 2450 2455 2460
 Gln Ala Leu Gly Met Met Asn Val Asp Gly Glu Leu Thr Ala Lys Glu
 2465 2470 2475 2480
 Leu Glu Lys Leu Lys Arg Ile Ile Asp Lys Leu Gln Gly Leu Thr Lys
 2485 2490 2495
 Glu Gln Cys Leu Asn Cys
 2500

<210> 2

<211> 1457

<212> PRT

<213> Porcine reproductive and respiratory syndrome virus

<400> 2

Leu Ala Ala Ser Asp Leu Thr Arg Cys Gly Arg Gly Gly Leu Val Val

1	5	10	15
Thr Glu Thr	Ala Val Lys Ile Val	Lys Phe His Asn Arg	Thr Phe Thr
	20	25	30
Leu Gly Pro	Val Asn Leu Lys	Val Ala Ser Glu	Val Glu Leu Lys Asp
	35	40	45
Ala Val Glu	His Asn Gln His	Pro Val Ala Arg	Pro Ile Asp Gly Gly
	50	55	60
Val Val Leu	Leu Arg Ser Ala	Val Pro Ser Leu	Ile Asp Val Leu Ile
	65	70	75
Ser Gly Ala	Asp Ala Ser Pro	Lys Leu Leu Ala	His His Gly Pro Gly
	85	90	95
Asn Thr Gly	Ile Asp Gly Thr	Leu Trp Asp Phe	Glu Ser Glu Ala Thr
	100	105	110
Lys Glu Glu	Val Ala Leu Ser	Ala Gln Ile Ile	Gln Ala Cys Asp Ile
	115	120	125
Arg Arg Gly	Asp Ala Pro Glu	Ile Gly Leu Pro	Tyr Lys Leu Tyr Pro
	130	135	140
Val Arg Gly	Asn Pro Glu Arg	Val Lys Gly Val	Leu Gln Asn Thr Arg
	145	150	155
Phe Gly Asp	Ile Pro Tyr Lys	Thr Pro Ser Asp	Thr Gly Ser Pro Val
	165	170	175
His Ala Ala	Ala Cys Leu Thr	Pro Asn Ala Thr	Pro Val Thr Asp Gly
	180	185	190
Arg Ser Val	Leu Ala Thr Thr	Met Pro Pro Gly	Phe Glu Leu Tyr Val
	195	200	205
Pro Thr Ile	Pro Ala Ser Val	Leu Asp Tyr Leu	Asp Ser Arg Pro Asp
	210	215	220
Cys Pro Lys	Gln Leu Thr Glu	His Gly Cys Glu	Asp Ala Ala Leu Lys
	225	230	235
Asp Leu Ser	Lys Tyr Asp Leu	Ser Thr Gln Gly	Phe Val Leu Pro Gly
	245	250	255
Val Leu Arg	Leu Val Arg Lys	Tyr Leu Phe Ala	His Val Gly Lys Cys
	260	265	270
Pro Pro Val	His Arg Pro Ser	Thr Tyr Pro Ala	Lys Asn Ser Met Ala
	275	280	285
Gly Ile Asn	Gly Asn Arg Phe	Pro Thr Lys Asp	Ile Gln Ser Val Pro
	290	295	300
Glu Ile Asp	Val Leu Cys Ala	Gln Ala Val Arg	Glu Asn Trp Gln Thr

305					310					315					320
Val	Thr	Pro	Cys	Thr	Leu	Lys	Lys	Gln	Tyr	Cys	Gly	Lys	Lys	Lys	Thr
				325					330					335	
Arg	Thr	Ile	Leu	Gly	Thr	Asn	Asn	Phe	Ile	Ala	Leu	Ala	His	Arg	Ala
			340					345					350		
Val	Leu	Ser	Gly	Val	Thr	Gln	Gly	Phe	Met	Lys	Lys	Ala	Phe	Asn	Ser
		355					360					365			
Pro	Ile	Ala	Leu	Gly	Lys	Asn	Lys	Phe	Lys	Glu	Leu	Gln	Thr	Pro	Val
	370					375					380				
Leu	Gly	Arg	Cys	Leu	Glu	Ala	Asp	Leu	Ala	Ser	Cys	Asp	Arg	Ser	Thr
385					390					395					400
Pro	Ala	Ile	Val	Arg	Trp	Phe	Ala	Ala	Asn	Leu	Leu	Tyr	Glu	Leu	Ala
				405					410					415	
Cys	Ala	Glu	Glu	His	Leu	Pro	Ser	Tyr	Val	Leu	Asn	Cys	Cys	His	Asp
			420					425					430		
Leu	Leu	Val	Thr	Gln	Ser	Gly	Ala	Val	Thr	Lys	Arg	Gly	Gly	Leu	Ser
		435					440					445			
Ser	Gly	Asp	Pro	Ile	Thr	Ser	Val	Ser	Asn	Thr	Ile	Tyr	Ser	Leu	Val
	450					455					460				
Ile	Tyr	Ala	Gln	His	Met	Val	Leu	Ser	Tyr	Phe	Lys	Ser	Gly	His	Pro
465					470					475					480
His	Gly	Leu	Leu	Phe	Leu	Gln	Asp	Gln	Leu	Lys	Phe	Glu	Asp	Met	Leu
				485					490					495	
Lys	Val	Gln	Pro	Leu	Ile	Val	Tyr	Ser	Asp	Asp	Leu	Val	Leu	Tyr	Ala
			500					505					510		
Glu	Ser	Pro	Thr	Met	Pro	Asn	Tyr	His	Trp	Trp	Val	Glu	His	Leu	Asn
		515					520					525			
Leu	Met	Leu	Gly	Phe	Gln	Thr	Asp	Pro	Lys	Lys	Thr	Ala	Ile	Thr	Asp
	530					535					540				
Ser	Pro	Ser	Phe	Leu	Gly	Cys	Arg	Ile	Ile	Asn	Gly	Arg	Gln	Leu	Val
545					550					555					560
Pro	Asn	Arg	Asp	Arg	Ile	Leu	Ala	Ala	Leu	Ala	Tyr	His	Met	Lys	Ala
				565					570					575	
Ser	Asn	Val	Ser	Glu	Tyr	Tyr	Ala	Ser	Ala	Ala	Ala	Ile	Leu	Met	Asp
			580					585					590		
Ser	Cys	Ala	Cys	Leu	Glu	Tyr	Asp	Pro	Glu	Trp	Phe	Glu	Glu	Leu	Val
		595					600					605			
Val	Gly	Ile	Ala	Gln	Cys	Ala	Arg	Lys	Asp	Gly	Tyr	Ser	Phe	Pro	Gly

610					615					620					
Thr 625	Pro	Phe	Phe	Met	Ser 630	Met	Trp	Glu	Lys	Leu 635	Arg	Ser	Asn	Tyr	Glu 640
Gly	Lys	Lys	Ser	Arg 645	Val	Cys	Gly	Tyr	Cys 650	Gly	Ala	Pro	Ala	Pro 655	Tyr
Ala	Thr	Ala	Cys 660	Gly	Leu	Asp	Val	Cys 665	Ile	Tyr	His	Thr	His 670	Phe	His
Gln	His	Cys 675	Pro	Val	Thr	Ile	Trp 680	Cys	Gly	His	Pro	Ala 685	Gly	Ser	Gly
Ser 690	Cys	Ser	Glu	Cys	Lys	Ser 695	Pro	Val	Gly	Lys	Gly 700	Thr	Ser	Pro	Leu
Asp 705	Glu	Val	Leu	Glu	Gln 710	Val	Pro	Tyr	Lys	Pro 715	Pro	Arg	Thr	Val	Ile 720
Met	His	Val	Glu	Gln 725	Gly	Leu	Thr	Pro	Leu 730	Asp	Pro	Gly	Arg	Tyr 735	Gln
Thr	Arg	Arg	Gly 740	Leu	Val	Ser	Val	Arg 745	Arg	Gly	Ile	Arg	Gly 750	Asn	Glu
Val	Gly	Leu 755	Pro	Asp	Gly	Asp	Tyr 760	Ala	Ser	Thr	Ala	Leu 765	Leu	Pro	Thr
Cys 770	Lys	Glu	Ile	Asn	Met	Val	Ala 775	Val	Ala	Ser	Asn 780	Val	Leu	Arg	Ser
Arg 785	Phe	Ile	Ile	Gly	Pro 790	Pro	Gly	Ala	Gly	Lys 795	Thr	Tyr	Trp	Leu 800	Leu
Gln	Gln	Val	Gln	Asp 805	Gly	Asp	Val	Ile	Tyr 810	Thr	Pro	Thr	His	Gln 815	Thr
Met	Leu	Asp	Met 820	Ile	Arg	Ala	Leu	Gly 825	Thr	Cys	Arg	Phe	Asn 830	Val	Pro
Ala	Gly	Thr 835	Thr	Leu	Gln	Phe	Pro 840	Val	Pro	Ser	Arg	Thr 845	Gly	Pro	Trp
Val 850	Arg	Ile	Leu	Ala	Gly	Gly 855	Trp	Cys	Pro	Gly	Lys 860	Asn	Ser	Phe	Leu
Asp 865	Glu	Ala	Ala	Tyr	Cys 870	Asn	His	Leu	Asp	Val 875	Leu	Arg	Leu	Leu	Ser 880
Lys	Thr	Thr	Leu	Thr 885	Cys	Leu	Gly	Asp	Phe 890	Lys	Gln	Leu	His	Pro 895	Val
Gly	Phe	Asp	Ser 900	His	Cys	Tyr	Val	Phe 905	Asp	Ile	Met	Pro	Gln 910	Thr	Gln
Leu	Lys	Thr	Ile	Trp	Arg	Phe	Gly	Gln	Asn	Ile	Cys	Asp	Ala	Ile	Gln

915					920					925						
Pro	Asp	Tyr	Arg	Asp	Lys	Leu	Met	Ser	Met	Val	Asn	Thr	Thr	Arg	Val	
930					935					940						
Thr	Tyr	Val	Glu	Lys	Pro	Val	Arg	Tyr	Gly	Gln	Val	Leu	Thr	Pro	Tyr	
945					950					955					960	
His	Arg	Asp	Arg	Glu	Asp	Asp	Ala	Ile	Thr	Ile	Asp	Ser	Ser	Gln	Gly	
965					970					975						
Ala	Thr	Phe	Asp	Val	Val	Thr	Leu	His	Leu	Pro	Thr	Lys	Asp	Ser	Leu	
980					985					990						
Asn	Arg	Gln	Arg	Ala	Leu	Val	Ala	Ile	Thr	Arg	Ala	Arg	His	Ala	Ile	
995					1000					1005						
Phe	Val	Tyr	Asp	Pro	His	Arg	Gln	Leu	Gln	Gly	Leu	Phe	Asp	Leu	Pro	
1010					1015					1020						
Ala	Lys	Gly	Thr	Pro	Val	Asn	Leu	Ala	Val	His	Cys	Asp	Gly	Gln	Leu	
1025					1030					1035					1040	
Ile	Val	Leu	Asp	Arg	Asn	Asn	Lys	Glu	Cys	Thr	Val	Ala	Gln	Ala	Leu	
1045					1050					1055						
Gly	Asn	Gly	Asp	Lys	Phe	Arg	Ala	Thr	Asp	Lys	Arg	Val	Val	Asp	Ser	
1060					1065					1070						
Leu	Arg	Ala	Ile	Cys	Ala	Asp	Leu	Glu	Gly	Ser	Ser	Ser	Pro	Leu	Pro	
1075					1080					1085						
Lys	Val	Ala	His	Asn	Leu	Gly	Phe	Tyr	Phe	Ser	Pro	Asp	Leu	Thr	Gln	
1090					1095					1100						
Phe	Ala	Lys	Leu	Pro	Val	Glu	Leu	Ala	Pro	His	Trp	Pro	Val	Val	Ser	
1105					1110					1115					1120	
Thr	Gln	Asn	Asn	Glu	Lys	Trp	Pro	Asp	Arg	Leu	Val	Ala	Ser	Leu	Arg	
1125					1130					1135						
Pro	Ile	His	Lys	Tyr	Ser	Arg	Ala	Cys	Ile	Gly	Ala	Gly	Tyr	Met	Val	
1140					1145					1150						
Gly	Pro	Ser	Val	Phe	Leu	Gly	Thr	Pro	Gly	Val	Val	Ser	Tyr	Tyr	Leu	
1155					1160					1165						
Thr	Lys	Phe	Val	Lys	Gly	Gly	Ala	Gln	Val	Leu	Pro	Glu	Thr	Val	Phe	
1170					1175					1180						
Ser	Thr	Gly	Arg	Ile	Glu	Val	Asp	Cys	Arg	Glu	Tyr	Leu	Asp	Asp	Arg	
1185					1190					1195					1200	
Glu	Arg	Glu	Val	Ala	Ala	Ser	Leu	Pro	His	Gly	Phe	Ile	Gly	Asp	Val	
1205					1210					1215						
Lys	Gly	Thr	Thr	Val	Gly	Gly	Cys	His	His	Val	Thr	Ser	Arg	Tyr	Leu	

1220	1225	1230
Pro Arg Val Leu Pro Lys Glu Ser Val Ala Val Val Gly Val Ser Ser 1235 1240 1245		
Pro Gly Lys Ala Ala Lys Ala Leu Cys Thr Leu Thr Asp Val Tyr Leu 1250 1255 1260		
Pro Asp Leu Glu Ala Tyr Leu His Pro Glu Thr Gln Ser Lys Cys Trp 1265 1270 1275 1280		
Lys Met Met Leu Asp Phe Lys Glu Val Arg Leu Met Val Trp Lys Asp 1285 1290 1295		
Lys Thr Ala Tyr Phe Gln Leu Glu Gly Arg Tyr Phe Thr Trp Tyr Gln 1300 1305 1310		
Leu Ala Ser Tyr Ala Ser Tyr Ile Arg Val Pro Val Asn Ser Thr Val 1315 1320 1325		
Tyr Leu Asp Pro Cys Met Gly Pro Ala Leu Cys Asn Arg Arg Val Val 1330 1335 1340		
Gly Ser Thr His Trp Gly Ala Asp Leu Ala Val Thr Pro Tyr Asp Tyr 1345 1350 1355 1360		
Gly Ala Lys Ile Ile Leu Ser Ser Ala Tyr His Gly Glu Met Pro Pro 1365 1370 1375		
Gly Tyr Lys Ile Leu Ala Cys Ala Glu Phe Ser Leu Asp Asp Pro Val 1380 1385 1390		
Lys Tyr Lys His Thr Trp Gly Phe Glu Ser Asp Thr Ala Tyr Leu Tyr 1395 1400 1405		
Glu Phe Thr Gly Asn Gly Glu Asp Trp Glu Asp Tyr Asn Asp Ala Phe 1410 1415 1420		
Arg Ala Arg Gln Glu Gly Lys Ile Tyr Lys Ala Thr Ala Thr Ser Leu 1425 1430 1435 1440		
Lys Phe Tyr Phe Pro Pro Gly Pro Val Ile Glu Pro Thr Leu Gly Leu 1445 1450 1455		

Asn

<210> 3

<211> 256

<212> PRT

<213> Porcine reproductive and respiratory syndrome virus

<400> 3

Met Lys Trp Gly Pro Cys Lys Ala Phe Leu Thr Lys Leu Ala Asn Phe
1 5 10 15

Leu Trp Met Leu Ser Arg Ser Ser Trp Cys Pro Leu Leu Ile Ser Leu
 20 25 30
 Tyr Phe Trp Pro Phe Cys Leu Ala Ser Pro Ser Pro Val Gly Trp Trp
 35 40 45
 Ser Phe Ala Ser Asp Trp Phe Ala Pro Arg Tyr Ser Val Arg Ala Leu
 50 55 60
 Pro Phe Thr Leu Ser Asn Tyr Arg Arg Ser Tyr Glu Ala Phe Leu Ser
 65 70 75 80
 Gln Cys Gln Val Asp Ile Pro Thr Trp Gly Thr Lys His Pro Leu Gly
 85 90 95
 Met Leu Trp His His Lys Val Ser Thr Leu Ile Asp Glu Met Val Ser
 100 105 110
 Arg Arg Met Tyr Arg Ile Met Glu Lys Ala Gly Gln Ala Ala Trp Lys
 115 120 125
 Gln Val Val Ser Glu Ala Thr Leu Ser Arg Ile Ser Ser Leu Asp Val
 130 135 140
 Val Ala His Phe Gln His Leu Ala Ala Ile Glu Ala Glu Thr Cys Lys
 150 155 160
 Tyr Leu Ala Ser Arg Leu Pro Met Leu His Asn Leu Arg Met Thr Gly
 165 170 175
 Ser Asn Val Thr Ile Val Tyr Asn Ser Thr Leu Asn Gln Val Phe Ala
 180 185 190
 Ile Phe Pro Thr Pro Gly Ser Arg Pro Lys Leu His Asp Phe Gln Gln
 195 200 205
 Trp Leu Ile Ala Val His Ser Ser Ile Phe Ser Ser Val Ala Ala Ser
 210 215 220
 Cys Thr Leu Phe Val Val Leu Trp Leu Arg Val Pro Ile Leu Arg Thr
 225 230 235 240
 Val Phe Gly Phe Arg Trp Leu Gly Ala Ile Phe Leu Ser Asn Ser Gln
 245 250 255

<210> 4

<211> 7509

<212> DNA

<213> Porcine reproductive and respiratory syndrome virus

<400> 4

atgtctgga tacttgatcg gtgcacgtgt accccaatg ccagggtgtt tatggcggag 60
 ggccaagtct actgcacacg atgcctcagt gcacggtctc tccttcccct gaacctccaa 120

gtttctgagc	tgggggtgct	aggcctatct	tacaggcccg	aagagccact	ccggtggacg	180
ttgccacgtg	cattccccac	tgttgagtgc	tccccgcg	gggcctgctg	gctttctgca	240
atctttccaa	tgcacgaat	gaccagtggg	aacctgaact	tccaacaaag	aatggtacgg	300
gtcgcagctg	agctttacag	agccggccag	ctcaccctg	cagtcttgaa	ggctctacaa	360
gtttatgaac	ggggttgccg	ctggtacccc	attgttgagc	ctgtccctgg	agtggccggt	420
ttcgccaatt	ccctacatgt	gagtgataaa	cctttcccg	gagcaactca	cgtgttgacc	480
aacctgccgc	tcccgacag	acccaagcct	gaagactttt	gcccccttga	gtgtgctatg	540
gctactgtct	atgacattgg	tcatgacgcc	gtcatgtatg	tggccgaaag	gaaagtctcc	600
tgggccctc	gtggcgggga	tgaagtga	tttgaagctg	tccccgggga	gttgaagttg	660
attgcaaac	ggctccgcac	ctccttcccg	ccccaccaca	cagtggacat	gtctaagttc	720
gccttcacag	cccctgggtg	tgggtgtttc	atgcggtcg	aacgccaaca	cggctgcctt	780
cccgtgaca	ctgtccctga	aggcaactgc	tgggtggagc	tgtttgactt	gcttccactg	840
gaagttcaga	acaaagaaat	tgcctatgct	aaccaatttg	gctaccagac	caagcatggt	900
gtctctggca	agtacctaca	gcggaggctg	caagttaatg	gtctccgagc	agtaactgac	960
ctaaacggac	ctatcgtcgt	acagtacttc	tccgttaagg	agagttggat	ccgccatttg	1020
aaactggcgg	gagaaccag	ctactctggg	tttgaggacc	tcctcagaat	aagggttgag	1080
cctaacacgt	cgccattggc	tgacaaggaa	gaaaaaattt	tccggtttgg	cagtcacaag	1140
tggtagcggc	ctggaaagag	agcaagaaaa	gcacgctctt	gtgcgactgc	tacagtgcgt	1200
ggccgcgctt	tgtccgttcg	tgaaccgcg	caggccaagg	agcacgaggt	tgccggcgcc	1260
aacaaggctg	agcacctcaa	acactactcc	ccgcctgccg	aagggaattg	tggttggcac	1320
tgcatttccg	ccatcgccaa	ccggatgggtg	aattccaaat	ttgaaaccac	ccttcccgaa	1380
agagtgcagc	ctccagatga	ctgggctact	gacgaggatc	ttgtgaatgc	catccaaatc	1440
ctcagactcc	ctgcggcctt	agacaggaac	ggtgcttgta	ctagcgccaa	gtacgtactt	1500
aagctggaag	gtgagcattg	gactgtcact	gtgacccctg	ggatgtcccc	ttctttgctc	1560
cctcttgaat	gtgttcaggg	ctgttgtggg	cacaagggcg	gtcttgggtc	cccagatgca	1620
gtcgagggtc	ccgtgcttgc	ccctgcctgg	cttgaccggc	tggctgaggt	gatgcactg	1680
cctagcagtg	ctatcccagc	cgctctggcc	gaaatgtctg	gcgattccga	tcgttcggct	1740
tctccggtca	ccaccgtgtg	gactgtttcg	cagttctttg	cccgtcacag	cggagggaat	1800
caccctgacc	aagtgcgctt	agggaaaatt	atcagccttt	gtcaggtgat	tgaggactgc	1860
tgctgttccc	agaacaaaac	caaccgggtc	accccgagg	aggtcgcagc	aaagattgac	1920
ctgtacctcc	gtggtgcaac	aaatcttgaa	gaatgcttgg	ccaggcttga	gaaagcgcg	1980
ccgccacg	taatcgacac	ctcctttgat	tgggatgttg	tgctccctgg	ggttgaggcg	2040
gcaacccaga	cgatcaagct	gccccaggtc	aaccagtgtc	gtgctctggt	ccctgtttgt	2100
actcaaaagt	ccttggacaa	caactcggtc	ccccagacc	ccttttctact	ggctaactac	2160
tactaccgtg	cgcaaggtga	cgaagttcgt	accgtgaaa	gactaaccgc	cgtgctctcc	2220
aagttggaaa	aggttggttcg	agaagaatat	gggctcatgc	caaccgagcc	tgggtccacg	2280
cccacactgc	cacgcgggct	cgacgaactc	aaagaccaga	tggaggagga	cttgctgaaa	2340
ctggctaacg	cccagacgac	ttcggacatg	atggcctggg	cagtcgagca	ggttgacctc	2400
aaaacttggg	tcaagaacta	cccgcggtgg	acaccaccac	ccccccgcgc	aaaagttcag	2460
cctcgaaaaa	cgaagcctgt	caagagcttg	ccggagagaa	agcctgtccc	cgcgccgcgc	2520
aggaaggttg	ggtccgattg	tggcagcccc	gtttcattag	gcggcgatgt	ccctaacagt	2580
tgggaagatt	tggctgttag	tagccccctt	gatctcccga	ccccacctga	gccggcaaca	2640
ccttcaagtg	agctggtgat	tgtgtcctca	ccgcaatgca	tcttcaggcc	ggcgacaccc	2700
ttgagtgcgc	cggctccaat	tcccgacact	cgcggaactg	tgtctcgacc	ggtgacaccc	2760
ttgagtgcgc	cgatccctgt	gcccgcaccg	cggcgtaagt	ttcagcaggt	gaaaagattg	2820
agttcggcgg	cggcaatccc	accgtaccag	gacgagcccc	tggatttgtc	tgcttccctc	2880
cagactgaat	atgaggcctc	tccccagca	ccgcccagca	gcggggcggt	tctgggagta	2940
gaggggcatg	aagctgagga	aaccctgagt	gaaatctcgg	acatgtcggg	taacattaaa	3000
cctgcgtccg	tgatcatcaag	cagctccttg	tccagcgtga	gaatcacacg	cccaaaatac	3060
tcagctcaag	ccatcatcga	ctcgggcggg	ccctgcagtg	ggcatctcca	agaggtaaag	3120
gaaacatgcc	ttagtgtcat	gcgcgaggca	tgtgatgcga	ctaagcttga	tgacctgct	3180
acgcaggaat	ggctttctcg	catgtgggat	cgggtggaca	tgctgacttg	gcgcaacacg	3240
tctgtttacc	agggcatttg	caccttagat	ggcaggttaa	agttcctccc	aaaaatgata	3300
ctcgagacac	cgcgcacctc	tccgtgtgag	tttgtgatga	tgcttcacac	gcctgcacct	3360
tccgtaggtg	cggagagcga	ccttaccatt	ggctcagttg	ctactgaaga	tgttccacgc	3420
atcctcgaga	aaatagaaaa	tgtcggcgag	atggccaacc	agggaccctt	ggccttctcc	3480
gaggataaac	cggtagatga	ccaacttgtc	aacgaccccc	ggatatcgtc	gcggaggcct	3540

gacgagagca	catcagctcc	gtccgcaggg	acaggtggcg	cgggtcttt	taccgatttg	3600
ccgccttcag	atggcgcgga	tgcggacggg	ggggggccgt	ttcggacggg	aaaaagaaa	3660
gctgaaaggc	tctttgacca	actgagccgt	caggtttttg	acctcgtctc	ccatctccct	3720
gttttcttct	cacgcctttt	ctaccctggc	ggtggttatt	ctccgggtga	ttgggggttt	3780
gcagctttta	ctctattgtg	cctcttttta	tgttacagtt	accagacctt	tggatttgct	3840
ccccctcttg	gtgtgttttc	tgggtcttct	cggcgcggtc	gaatgggggt	ttttggtgc	3900
tggttggctt	ttgctgttgg	tctgttcaag	cctgtgtccg	accagtcggg	cgctgcttgt	3960
gagtttgact	cgccagagtg	tagaaacatc	cttcattctt	ttgagcttct	caaaccttgg	4020
gaccctgttc	gcagccttgt	tgtgggcccc	gtcgggtctcg	gtcttgccat	tcttggcagg	4080
ttactgggcg	gggcacgctg	catctggcac	tttttgett	ggcttggcat	tgttgacagc	4140
tgtatcttgg	ctggagctta	cgtgctttct	caaggtaggt	gtaaaaagt	ctggggatct	4200
tgtataagaa	ctgctcctaa	tgaggtcgct	tttaacgtgt	ttcctttcac	acgtgcgacc	4260
aggtcgtcac	ttatcgacct	gtgcgatcgg	ttttgtgcgc	caaaaggaat	ggacccatt	4320
ttctctgcc	ctgggtggcg	cgggtgctgg	gccggccgaa	gccccattga	gcaaccctct	4380
gaaaaaccca	tcgcgtttgc	ccaattggat	gaaaagaaga	ttacggctag	gactgtggtc	4440
gccagacctt	atgaccccaa	ccaagccgta	aagtgttgc	gggtattgca	gtcgggtggg	4500
cgatggtggc	taagcggtcc	caaaagtggg	caaggtttcc	gctgttccat	tccgagcccc	4560
ttctttccca	ctggagtga	agttgacctt	gattgcaggg	tcgtggttga	ccctgacact	4620
ttactgcag	ctctccggtc	tggctactcc	accacaaacc	tcgtccttgg	tgtaggggac	4680
tttgcccagc	tgaatggatt	aaaaatcagg	caaatttcca	agccttcagg	gggaggccca	4740
catctcatgg	ctgccctgca	tgttgcctgc	tcgatggctc	tgcacatgct	tgtctgggatt	4800
tatgtgactg	cgggtgggtt	ttgcggcacc	ggcaccaacg	accctggttg	cgctaaccgg	4860
tttgccgtcc	ctggctacgg	acctggctct	ctctgcacgt	ccaggttgtg	catttcccaa	4920
cacggcctta	ccctgccctt	gacagcactt	gtggcgggat	tcgggtattca	agaaattgcc	4980
ttggtcggtt	tgatttttgt	ttccatcgga	ggcatggctc	ataggttgag	ctgtaaggct	5040
gacatgctgt	gtgttttggc	tgcaattggc	agctatgttt	gggtacctct	tacctggttg	5100
ctttgtgtgt	ttccttgctg	gttgcgctgt	ttttctttgc	acccctcac	catcctatgg	5160
ttggtgtttt	tcttgatttc	tgtgaatatg	ccttcaggaa	tcttgcccat	ggtgttgttg	5220
gtttctcttt	ggcttcttgg	tcgttatact	aatgttgctg	gccttgtcac	cccctacgac	5280
attcatcatt	acaccagtgg	ccccgcgggt	gttgccgcct	tggctaccgc	accagatggg	5340
acctacttgg	ccgctgtccg	ccgcgctgcg	ttgactggcc	gcacatgct	gtttaccccg	5400
tcccagcttg	ggtctcttct	tgaggggtgt	ttcagaactc	gaaagccctc	actgaacacc	5460
gtcaatgtga	tcgggtccctc	catgggtctc	ggcgggggtg	ttaccatcga	cgggaaagtc	5520
aagtgcgtaa	ctgccgcaca	tgtccttacg	ggcaattcag	ctcgggtttc	cggggtcggc	5580
ttcaatcaaa	tgcttgactt	tgacgtaaag	ggagatttcc	ctatagctga	ttgccogaat	5640
tggcaagggg	ctgcccccaa	gacccaattc	tgcacggatg	gatggactgg	ccgtgcctat	5700
tggctaacat	cctctggcgt	cgaacccggc	gtcattggaa	aaggattcgc	cttctgcttc	5760
accgcatgtg	gcgattccgg	gtccccagtg	atcacaggag	ccggtgagct	tgtcggcggt	5820
cacacgggat	cgaataaaca	aggggggggc	attgttacgc	gcccctcagg	ccagttttgt	5880
aatgtggcac	ccatcaagct	aagcgaatta	agtgaattct	ttgctgggcc	taaggtcccg	5940
ctcgggtgat	tgaaggtcgg	cagccacata	attaaagaca	taagcgagggt	gccttcagat	6000
ctttgtgcct	tgcttgctgc	caaacctgaa	ctggaaggag	gcctctccac	cgtccaactt	6060
ctttgtgtgt	tttttctcct	gtggagaatg	atgggacatg	cctggacgcc	cttggttgct	6120
gtgagtttct	ttattttgaa	tgaggttctc	ccagccgtcc	tggtcgggag	tgttttctcc	6180
tttggaatgt	ttgtgctatc	ctggctcacg	ccatggtctg	cgcaagttct	gatgatcagg	6240
cttctgacag	cagctcttaa	caggaacaga	tggctcattg	cctttttcag	cctcggtgca	6300
gtgaccggtt	ttgtcgcaga	tcttgcggcc	actcaggggc	atccgttgca	ggcagtgatg	6360
aatttgagca	cctatgcatt	cctgcctcgg	atgatggttg	tgacctcacc	agtcccagtg	6420
atcacgtgtg	gtgtcgtgca	cctacttgcc	atcattttgt	acttgtttaa	gtaccgtggc	6480
ccgcaccata	tccttggttg	cgatggagt	ttctctgcgg	ctttcttctt	gagatacttt	6540
gccgagggaa	agttgagggg	aggggtgtcg	caatcctgcg	gaatgaatca	tgagtctctg	6600
actggtgccc	tcgctatgag	actcaatgac	gaggacttgg	atttccttat	gaaatggact	6660
gatttttaagt	gctttgtttc	tgcgtccaac	atgaggaatg	cagcgggtca	atttatcgag	6720
gctgcctatg	ctaaagcact	tagagtagaa	ctggcccagt	tgggtgcagg	tgataaagtt	6780
cgaggtactt	tggccaaact	tgaagctttt	gctgataacc	tggcacctca	actctcgccc	6840
ggtgacattg	ttgtcgtctc	cggccacacg	cctgttggca	gtatcttcga	cctaaagggt	6900
ggtagcacca	agcataacct	ccaagccatt	gagaccagag	tccttgctgg	gtccaaaatg	6960

accgtggcgc	gcgctcgtcga	cccgaccccc	acgccccccac	ccgcacccgt	gccccatcccc	7020
ctcccaccga	aagttcttga	gaatggcccc	aacgcttgagg	gggatgagga	ccgtttgaat	7080
aagaagaaga	ggcgaggat	ggaagccctc	ggcatctatg	ttatgggcgg	gaaaaagtac	7140
cagaaatttt	gggacaagaa	ttccggtgat	gtgttttatg	aggaggtcca	taataacaca	7200
gatgagtggg	agtgtctcag	agttggcgac	cctgccgact	ttgaccctga	gaaggggaact	7260
ctgtgtggac	atgtcaccat	tgaaaacaag	gcttaccatg	tttacacctc	cccatctggt	7320
aagaagtctt	tggtccccgt	caaccagag	aatggaagag	ttcaatggga	agctgcaaag	7380
ctttccgtgg	agcaggccct	aggtatgatg	aatgtcgacg	gcgaactgac	tgccaaagaa	7440
ctggagaaac	tgaaaagaat	aattgacaaa	ctccagggcc	tgactaagga	gcagtgttta	7500
aactgctag						7509

<210> 5

<211> 4374

<212> DNA

<213> Porcine reproductive and respiratory syndrome virus

<400> 5

ctagccgccca	gcgacttgac	ccgctgtggt	cgcgggcggt	tggttggttac	tgaaacagcg	60
gtaaaaatag	tcaaatttca	caaccggacc	ttcacccctgg	gacctgtgaa	tttaaaagtg	120
gccagtgagg	ttgagctaaa	agacgcgggt	gagcacaacc	aacacccggt	tgcgagaccg	180
atcgatggtg	gagttgtgct	cctgcgttcc	gcggttcctt	cgcttataga	cgtcttgatc	240
tccggtgctg	atgcatctcc	caagttactt	gccccatcac	ggccgggaaa	cactgggatc	300
gatggcacgc	tctgggattt	tgagtccgaa	gccactaaag	aggaagtcgc	actcagtgcg	360
caataataac	aggcttgtga	cattaggcgc	ggcgacgctc	ctgaaattgg	tctcccttac	420
aagctgtacc	ctgttagggg	taaccctgag	cggttgaaag	gagttctgca	gaatacaagg	480
tttgagagaca	taccttacia	aacccccagt	gacactggaa	gcccagtgcg	cgcggtgcc	540
tgccctacgc	ccaacgccac	tccggtgact	gatgggcgct	ccgtcttggc	cacgaccatg	600
ccccccgggt	ttgagttata	tgtaccgacc	ataccagcgt	ctgtccttga	ttaccttgac	660
tctaggcctg	actgccctaa	acagctgaca	gagcacggct	gcgaagatgc	cgactgaaa	720
gacctctcta	aatatgactt	gtccacccaa	ggctttgttt	tacctggagt	tcttcgcctt	780
gtgcggaaat	acctgtttgc	ccatgtaggt	aagtgccac	ccgttcacgc	gccttctact	840
tacctgtcta	agaattctat	ggctggaata	aatgggaaca	ggttcccaac	caaggacatt	900
cagagcgtcc	ctgaaatcga	cgctctgtgc	gcacaggctg	tgcgagaaaa	ctggcaaaact	960
gtcacccctt	gtactcttaa	gaaacagtat	tgcggggaaga	agaagactag	gaccatactc	1020
ggcaccaata	acttcatcgc	actagcccac	cgagcagtg	tgagtgggtg	taccaggggc	1080
ttcatgaaaa	aggcgtttaa	ctcgcccatc	gccctcgga	agaacaagtt	taaggagcta	1140
cagactccgg	tctggggcag	gtgccttgaa	gctgatctcg	catcctgcga	tcgatccacg	1200
cctgcaattg	tccgctgggt	tgccgccaac	cttctttatg	aacttgccctg	tgctgaagag	1260
catctaccgt	cgtacgtgct	gaactgctgc	cacgacttac	tggtcacgca	gtccggcgca	1320
gtgactaaga	gaggtggcct	gtcgtctggc	gacccgatca	cctctgtgtc	taacaccatt	1380
tatagtttgg	tgatctatgc	acagcatatg	gtgcttagtt	acttcaaaaag	tggtcacccc	1440
catggccttc	tgttcttaca	agaccagcta	aagtttgagg	acatgctcaa	ggttcaacct	1500
ctgatcgtct	attcggacga	cctcgtgctg	tatgccgagt	ctccaccat	gccaaactat	1560
cactgggtggg	ttgaacatct	gaatttgatg	ctgggggttc	agacggaccc	aaagaagaca	1620
gcaataacag	actcgccatc	atctcttaggc	tgtagaataa	taaattggcg	ccagctagtc	1680
cccaaccgtg	acaggatcct	cgcgccctc	gcctatcaca	tgaaggcgag	taatgtttct	1740
gaataactatg	cctcagcggc	tgcaatactc	atggacagct	gtgcttggtt	ggagtatgat	1800
cctgaatggg	ttgaagaact	tgtagttgga	atagcgagct	gcgcccgcga	ggacggctac	1860
agctttcccg	gcacgcggtt	cttcatgtcc	atgtgggaaa	aactcaggtc	caattatgag	1920
gggaagaagt	cgagagtgtg	cgggtactgc	ggggcccg	ccccgtacgc	tactgctgtg	1980
ggcctcgacg	tctgcattta	ccacaccac	ttccaccagc	attgtccagt	cacaatctgg	2040
tgtggccatc	cagcgggttc	tggttcttgt	agttagtgca	aatcccctgt	agggaaaggc	2100
acaagccctt	tagacgaggt	gctggaacaa	gtcccgtata	agccccacg	gaccgttatc	2160
atgcatgtgg	agcagggtct	cacccccctt	gatccaggta	gataccaaac	tcgcccgcga	2220
ttagtctctg	tcaggcgtgg	aattagggga	aatgaagttg	gactaccaga	cggtgattat	2280
gctagcaccg	ccttgctccc	tacctgcaaa	gagatcaaca	tggtcgctgt	cgcttccaat	2340

gtattgcgca	gcaggttcat	catcgGCCca	cccggTgctg	ggaaaacata	ctggctcctt	2400
caacaggTcc	aggatggTga	tgTtatTTac	acaccaactc	accagaccat	gcttgacatg	2460
attagggtctt	tggggacgtg	ccggttcaac	gtcccggcag	gcacaacgct	gcaattcccc	2520
gtccccctccc	gcaccggTcc	gtgggttcgc	atcctagccg	gcggttggtg	tcctggcaag	2580
aattccttcc	tagatgaagc	agcgtattgc	aatcaccttg	atgttttgag	gcttcttagt	2640
aaaactaccc	tcacctgtct	aggagacttc	aagcaactcc	acccagtggg	ttttgattct	2700
cattgctatg	tttttgacat	catgcctcaa	actcaactga	agaccatctg	gaggtttgga	2760
cagaatatct	gtgatgccat	tcagccagat	tacagggaca	aactcatgtc	catggtcaac	2820
acaaccctg	tgacctacgt	ggaaaaacct	gtcaggTatg	ggcaggTcct	cacccctac	2880
cacagggacc	gagaggacga	cgccatcact	attgactcca	gtcaaggcgc	cacattcgat	2940
gtgggttacat	tgcatTTgcc	cactaaagat	tactcaaca	ggcaaagagc	ccttgTtgct	3000
atcaccagg	caagacacgc	tatctttgtg	tatgaccac	acaggcagct	gcagggttg	3060
tttgatcttc	ctgcaaaaagg	cacgccgctc	aacctcgcag	tgactgcga	cgggcagctg	3120
atcgtgctgg	atagaaataa	caaagaatgc	acggttgctc	aggctctagg	caacggggat	3180
aaatttaggg	ccacagacaa	gcgtgttgta	gattctctcc	gcgccatttg	tgtgatcta	3240
gaagggtcga	gctctccgct	ccccaggTc	gcacacaact	tgggatttta	tttctcacct	3300
gatttaacac	agtttgctaa	actcccagta	gaacttgcac	ctcactggcc	cgtggtgtca	3360
accagaaca	atgaaaagt	gccggatcgg	ctggttgcca	gccttcgccc	tatccataaa	3420
tacagccg	cgtgcatcgg	tgccggctat	atggtggg	cttcggtgtt	tctaggcact	3480
cctggggctg	tgtcacta	tctcaca	tttgTtaagg	gcggggctca	agtgttccg	3540
gagacggTtt	tcagcaccgg	ccgaattgag	gtagactgcc	gggaatatct	tgatgatcgg	3600
gagcgagaag	ttgctgcgtc	cctcccacac	ggtttcattg	gcgacgtcaa	aggcactacc	3660
gttgaggat	gtcatcatgt	cacctccaga	tacctccgc	gcgtccttc	caaggaatca	3720
gttgcggtag	tcggggTttc	aagccccgga	aaagccgcga	aagcattgtg	cacactgaca	3780
gatgtgtacc	tcccagatct	tgaagcctat	ctccaccgg	agaccagtc	caagtgtgg	3840
aaaatgatgt	tggagtTcaa	agaagttcga	ctaattgtct	ggaaagacaa	aacagcctat	3900
ttccaacttg	aaggtcgcta	tttcacctgg	tatcagcttg	ccagctatgc	ctcgtacatc	3960
cgtgttcccg	tcaactctac	ggtgtacttg	gaccttgca	tgggccccgc	cctttgcaac	4020
aggagagtgc	tcgggtccac	ccactggggg	gctgacctcg	cggTcacccc	ttatgattac	4080
ggcgctaaaa	ttatcctgtc	tagcgcgTac	catggtgaaa	tgccccccgg	atacaaaatt	4140
ctggcgTgcg	cggagtTctc	gttggaTgac	ccagTtaagt	acaaacatac	ctgggggttt	4200
gaatcggata	cagcgTatct	gtatgagTtc	accggaaacg	gtgaggactg	ggaggattac	4260
aatgatgcgt	ttcgtgcgcg	ccaggaagg	aaaatttata	aggccactgc	caccagcttg	4320
aagttttatt	ttcccccg	ccctgtcatt	gaaccaactt	taggcctgaa	ttga	4374

<210> 6

<211> 771

<212> DNA

<213> Porcine reproductive and respiratory syndrome virus

<400> 6

atgaaatggg	gtccatgcaa	agcctttttg	acaaaattgg	ccaacttttt	gtggatgctt	60
tcacggagtt	cttggtgtcc	attgttgata	tcattatatt	tttggccatt	ttgtttggct	120
tcaccatcgc	cggttggctg	gtggtctttt	gcacagatt	ggtttgctcc	gcgatactcc	180
gtacgcgcc	tgccattcac	tctgagcaat	tacagaagat	cttatgaggc	ctttctttcc	240
cagtccaag	tggacattcc	cacctgggga	actaaacatc	ctttggggat	gctttggcac	300
cataaggtgt	caaccctgat	tgatgaaatg	gtgtcgcgtc	gaatgtaccg	catcatggaa	360
aaagcagggc	aggctgcctg	gaaacaggTg	gtgagcgagg	ctacgctgtc	tcgcattagt	420
agtttggtatg	tggTggctca	ttttcagcat	ctagccgcca	ttgaagccga	gacctgtaaa	480
tatttgccct	cccggctgcc	catgctacac	aacctgcgca	tgacagggtc	aaatgtaacc	540
atagtgtata	atagcacttt	gaatcaggTg	tttgctattt	ttccaacccc	tggTtcccg	600
ccaaagcttc	atgattttca	gcaatggTta	atagctgtac	attcctccat	attttcctct	660
gttgacgctt	cttgtaactct	ttttgttgTg	ctgtggtTgc	gggttccaat	actacgtact	720
gtttttgggtt	tccgctgggtt	aggggcaatt	tttctttcga	actcacagt	a	771